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TECH CENTER 1600/2900

SEQUENCE LISTING

<110> Schweighoffer, Fabien  
Bracco, Laurent  
Tocque, Bruno

<120> Qualitative Differential Screening

<130> 50146/004002

<140> 09/623,828

<141> 2000-11-30

<150> PCT/FR99/00547

<151> 1999-03-11

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Gly Gln Leu Gly Gly Glu Glu Trp Thr Arg His Gly Ser Phe Val Asn  
20 25 30  
Lys Pro Thr Arg Gly Trp Leu His Pro Asn Asp Lys Val Met Gly Pro  
35 40 45  
Gly Val Ser Tyr Leu Val Arg Tyr Met Gly Cys Val Glu Val Leu Gln  
50 55 60  
Ser Met Arg Ala Leu Asp Phe Asn Thr Arg Thr Gln Val Thr Arg Glu  
65 70 75 80  
Ala Ile Ser Leu Val Cys Glu Ala Val Pro Gly Ala Lys Gly Ala Thr  
85 90 95  
Arg Arg Arg Lys Pro Cys Ser Arg Pro Leu Ser Ser Ile Leu Gly Arg  
100 105 110  
Ser Asn Leu Lys Phe Ala Gly Met Pro Ile Thr Leu Thr Val Ser Thr  
115 120 125  
Ser Ser Leu Asn Leu Met Ala Ala Asp Cys Lys Gln Ile Ile Ala Asn  
130 135 140  
His His Met Gln Ser Ile Ser Phe Ala Ser Gly Asp Pro Asp Thr  
145 150 155 160  
Ala Glu Tyr Val Ala Tyr Val Ala Lys Asp Pro Val Asn Gln Arg Ala  
165 170 175  
Cys His Ile Leu Glu Cys Pro Glu Gly Leu Ala Gln Asp Val Ile Ser  
180 185 190  
Thr Ile Gly Gln Ala Phe Glu Leu Arg Phe Lys Gln Tyr Leu Arg Asn  
195 200 205  
Pro Pro Lys Leu Val Thr Pro His Asp Arg Met Ala Gly Phe Asp Gly  
210 215 220  
Ser Ala Trp Asp Glu Glu Glu Glu Glu Pro Pro Asp His Gln Tyr Tyr  
225 230 235 240  
Asn Asp Phe Pro Gly Lys Glu Pro Pro Leu Gly Gly Val Val Asp Met  
245 250 255  
Arg Leu Arg Glu Gly Ala Ala Pro Gly Ala Ala Arg Pro Thr Ala Pro  
260 265 270  
Asn Ala Gln Thr Pro Ser His Leu Gly Ala Thr Leu Pro Val Gly Gln  
275 280 285  
Pro Val Gly Gly Asp Pro Glu Val Arg Lys Gln Met Pro Pro Pro Pro  
290 295 300  
Pro Cys Pro Gly Arg Glu Leu Phe Asp Asp Pro Ser Tyr Val Asn Val  
305 310 315 320  
Gln Asn Leu Asp Lys Ala Arg Gln Ala Val Gly Gly Ala Gly Pro Pro  
325 330 335  
Asn Pro Ala Ile Asn Gly Ser Ala Pro Arg Asp Leu Phe Asp Met Lys  
340 345 350  
Pro Phe Glu Asp Ala Leu Arg Val Pro Pro Pro Pro Gln Ser Val Ser  
355 360 365

Met	Ala	Glu	Gln	Leu	Arg	Gly	Glu	Pro	Trp	Phe	His	Gly	Lys	Leu	Ser
370						375					380				
Arg	Arg	Glu	Ala	Glu	Ala	Leu	Leu	Gln	Leu	Asn	Gly	Asp	Phe	Leu	Val
385					390					395					400
Arg	Thr	Lys	Asp	His	Arg	Phe	Glu	Ser	Val	Ser	His	Leu	Ile	Ser	Tyr
			405						410					415	
His	Met	Asp	Asn	His	Leu	Pro	Ile	Ile	Ser	Ala	Gly	Ser	Glu	Leu	Cys
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 cccaacgaca aagtcatggg acccgggggt tcctacttgg ttcggtacat gggttgtgtg 180  
 gaggtcctcc agtcaatgcg tgccctggac ttcaacaccc ggactcaggt caccagggag 240  
 gccatcagtc tgggtgtgtga ggctgtgccc ggtgctaagg gggcgacaag gaggagaaag 300  
 ccctgtagcc gcccgtcag ctctatcctg gggaggagta acctgaaatt tgctggaatg 360  
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 atcatcgcca accaccacat gcaatctatc tcatttgcac cggcggggga tccggacaca 480  
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 gagtgtcccg aagggcttgc ccaggatgtc atcagcacca ttggccaggc cttcgagttg 600  
 cgcttcaaac aatacctcag gaaccacccc aaactgggtc cccctcatga caggatggct 660  
 ggctttgatg gctcagcatg ggatgaggag gaggaagagc cacctgacca tcagtactat 720  
 aatgacttcc cggggaagga accccccttg gggggggtgg tagacatgag gcttcgggaa 780  
 ggagccgctc caggggctgc tcgacccact gcacccaatg cccagacccc cagccacttg 840  
 ggagctacat tgctgtagg acagcctgtt gggggagatc cagaagtccg caaacagatg 900  
 ccacctccag caccctgtcc aggcagagag ctttttgatg atccctccta tgtcaacgtg 960  
 cagaacctag acaaggcccc gcaagcagtg ggtggtgctg ggccccccaa tcctgctatc 1020  
 aatggcagtg caccgccgga cctgtttgac atgaagccct tcgaagatgc tcttcgggtg 1080  
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 gggaagctga gccggcgga ggctgaggca ctgctgcagc tcaatgggga cttcttggtt 1200  
 cggactaagg atcaccgctt tgaaagtgtc agtcacctta tcagctacca catggacaat 1260  
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and.